

p1075-seq-pc

## Sequence listing

&lt;110&gt; Epigenomics AG

<120> Method for investigating cytosine methylation in DNA by means of  
DNA repair enzymes

&lt;130&gt; 82325

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&lt;211&gt; 26

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; chemically treated genomic DNA (Homo sapiens)

&lt;400&gt; 1

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26

&lt;210&gt; 2

&lt;211&gt; 19

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; chemically treated genomic DNA (Homo sapiens)

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&lt;210&gt; 3

&lt;211&gt; 19

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; chemically treated genomic DNA (Homo sapiens)

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&lt;211&gt; 20

&lt;212&gt; DNA

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&lt;223&gt; chemically treated genomic DNA (Homo sapiens)

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&lt;211&gt; 20

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

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